

GenCore version 4.5
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OM protein - protein search, using sw model

Run on: September 4, 2002, 16:10:56 : Search time 99.82 Seconds
(without alignments)
53.833 Million cell updates/sec

Title: US-09-052-089A-3
Perfect score: 1066
Sequence: 1 RTINKLFPDLAEEENVLD.....DLQSDAKREIMSLKKLTMLQ 220

Scoring table:
BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 231628 seqs, 24425594 residues

Total number of hits satisfying chosen parameters: 231628

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :
Issued_Patents_AA:*
1: /cgn2.6/prodata/2/iaa/5A.COMB.pep:*
2: /cgn2.6/prodata/2/iaa/5B.COMB.pep:*
3: /cgn2.6/prodata/2/iaa/6A.COMB.pep:*
4: /cgn2.6/prodata/2/iaa/6B.COMB.pep:*
5: /cgn2.6/prodata/2/iaa/PCTUS.COMB.pep:*
6: /cgn2.6/prodata/2/iaa/backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	1066	100.0	220	4	US-09-052-089A-3
2	1066	100.0	469	4	US-09-052-089A-1
3	1046	98.1	469	2	US-08-968-751-2
4	906	85.0	220	4	US-09-052-089A-4
5	906	85.0	470	4	US-09-052-089A-2
6	156	14.6	2482	1	US-08-328-254-6
7	156	14.6	3248	1	US-08-333-700-1
8	156	14.6	3248	5	PCT-US95-16216-1
9	148.5	13.9	443	2	US-08-795-475-6
10	144	13.5	414	5	PCT-US93-03077-3
11	144	13.5	976	4	US-09-104-324B-4
12	144	13.5	1093	5	PCT-US93-03077-1
13	142	13.3	1090	4	US-09-085-199B-5
14	141	13.1	914	4	US-09-085-199B-4
15	140	13.1	756	4	US-09-085-199B-9
16	139	13.0	576	2	US-08-533-306A-2
17	139	13.0	576	2	US-08-742-923A-2
18	139	13.0	816	2	US-08-533-306A-6
19	139	13.0	816	2	US-08-742-923A-6
20	139	13.0	885	2	US-08-533-306A-4
21	139	13.0	885	2	US-08-742-923A-4
22	138.5	13.0	1939	4	US-09-310-187A-1
23	136.5	12.8	896	1	US-08-095-737-2
24	136.5	12.8	896	1	US-08-460-145-2
25	136.5	12.8	896	2	US-08-477-389-2
26	135.5	12.7	316	4	US-08-098-327E-31
27	135.5	12.7	316	4	US-08-462-625-31

28	133.5	12.5	386	4	US-09-085-199B-2	Sequence 2, Appl
29	132.5	12.4	376	6	5180810-1	Patent No. 5180810
30	132.5	12.4	1388	4	US-09-572-191-2	Sequence 2, Appl
31	132.5	12.4	1886	4	US-08-938-105-3	Sequence 3, Appl
32	130	12.2	2101	1	US-08-466-390-4	Sequence 4, Appl
33	130	12.2	2101	1	US-08-470-950-4	Sequence 4, Appl
34	130	12.2	2101	1	US-08-467-781-4	Sequence 4, Appl
35	130	12.2	2101	1	US-08-195-487-4	Sequence 4, Appl
36	130	12.2	2101	2	US-08-483-924-4	Sequence 4, Appl
37	130	12.2	2101	4	US-09-452-294-1	Sequence 1, Appl
38	130	12.2	2101	5	PCT-US93-06160-4	Sequence 4, Appl
39	129.5	12.1	351	1	US-08-402-217A-2	Sequence 2, Appl
40	129.5	12.1	351	1	US-08-700-178-2	Sequence 2, Appl
41	129.5	12.1	351	3	US-08-995-654-2	Sequence 2, Appl
42	128.5	12.1	1388	2	US-08-685-576-1	Sequence 1, Appl
43	128	12.0	887	1	US-08-095-737-4	Sequence 4, Appl
44	128	12.0	897	1	US-08-480-145-4	Sequence 4, Appl
45	128	12.0	897	2	US-08-477-389-4	Sequence 4, Appl

ALIGNMENTS

RESULT 1
US-09-052-089A-3
Sequence 3, Application US/09052089A
Patent No. 6346605
GENERAL INFORMATION:
APPLICANT: Lee, Soo Y.
Choi, Yongwon
TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNF RECEPTOR SUPER FAMILY, AND USGS THEREOF
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.
STREET: 411 Hackensack Ave, Continental Plaza, 4th Floor
CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
ZIP: 07601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/052,089A
FILING DATE: 31-Mar-1998
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
TELEFAX: 201-343-1684
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 220 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHEICAL: NO
FRAGMENT TYPE: Internal
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-052-089A-3
Query Match 100.0%; Score 1066; DB 4; Length 220;

Best Local Similarity 100.0%; Pred. No. 4.7e-85;
Matches 220; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 60
Db 1 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 60
OY 61 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 120
Db 61 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 120
OY 121 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 180
Db 121 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 180
OY 181 LQTVYSELDAQKLELSAQKDLQSDAKREIMSLKKLTMLQ 220
Db 181 LQTVYSELDAQKLELSAQKDLQSDAKREIMSLKKLTMLQ 220

RESULT 2

US-09-052-089a-1
Sequence 1, Application US/09052089a
Patent No. 6346605

GENERAL INFORMATION:

APPLICANT: Lee, Soo Y.

Choi, Yongwon

TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNF RECEPTOR SUPER

NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:

ADDRESSEE: David A. Jackson, Esq.

STREET: 411 Hackensack Ave, Continental Plaza, 4th

Floor

CITY: Hackensack

STATE: New Jersey

COUNTRY: USA

ZIP: 07601

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/052, 089a

FILING DATE: 31-Mar-1998

CLASSIFICATION: <Unknown>

ATTORNEY/AGENT INFORMATION:

NAME: Jackson Esq., David A.

REGISTRATION NUMBER: 26,742

REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1

TELECOMMUNICATION INFORMATION:

TELEPHONE: 201-487-5800

TELEFAX: 201-343-1684

INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:

LENGTH: 469 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

HYPOTHETICAL: NO

FRAGMENT TYPE: <Unknown>

ORIGINAL SOURCE:

ORGANISM: Homo sapiens

SEQUENCE DESCRIPTION: SEQ ID NO: 1:

US-09-052-089a-1

Query Match 100.0%; Score 1066; DB 4; Length 469;
Best Local Similarity 100.0%; Pred. No. 1.2e-84;
Matches 220; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 60
Db 56 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 115

OY 61 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 120
Db 116 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 175

OY 121 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 180
Db 176 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 235

OY 181 LQTVYSELDAQKLELSAQKDLQSDAKREIMSLKKLTMLQ 220
Db 226 LQTVYSELDAQKLELSAQKDLQSDAKREIMSLKKLTMLQ 275

RESULT 3

US-08-968-751-2
Sequence 2, Application US/08968751
Patent No. 5948643

GENERAL INFORMATION:

APPLICANT: Rubinfeld, Bonnie

Polakis, Paul G.

APPLICANT: Ligenfelter, Carol

APPLICANT: Vuong, Terilyn T.

TITLE OF INVENTION: MODULATORS OF BRCA1 ACTIVITY

NUMBER OF SEQUENCES: 6

CORRESPONDENCE ADDRESS:

ADDRESSEE: ONYX Pharmaceuticals, Inc.

STREET: 3031 Research Drive

CITY: Richmond

STATE: CA

COUNTRY: USA

ZIP: 94806

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

OPERATING SYSTEM: IBM PC compatible

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/968, 751

FILING DATE:

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Giotta, Gregory

REGISTRATION NUMBER: 32,028

REFERENCE/DOCKET NUMBER: ONYX1024 GG

TELECOMMUNICATION INFORMATION:

TELEPHONE: (510) 262-8710

TELEFAX: (510) 222-9758

INFORMATION FOR SEQ ID NO: 2:

SEQUENCE CHARACTERISTICS:

LENGTH: 469 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-968-751-2

Query Match 98.1%; Score 1046; DB 2; Length 469;
Best Local Similarity 98.6%; Pred. No. 6.5e-83;
Matches 217; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

OY 1 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 60
Db 56 RTIINKLFFDLAEEENVLDRFLKNELDNVRAQLSQKDKERDSQVITDRLRDTLEERN 115

OY 61 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 120
Db 116 ATTVSLQALGKAEMLCSTLKKOMKYLEQOODETKQAOEAGRLRSKKTMEQIELLLQS 175

OY 121 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 180
Db 121 QLPVEEEMIRMGVGSQSAVEOLAVYCVSLKKEYENLKARRASGEVADKLKRDLFSSRSK 180

Db 176 QREVEEMIRDMGVGQSAVEQLAVYCVSLKREYENLKREARRASGVADKLKRDLFSSRSK 235
QY 181 LQTVSELDQAKLELSAQKDLQSAKDEKIMSLKKTLMLQ 220
Db 236 LQTVSELDQAKLELSAQKDLQSAKDEKIMSLKKTLMLQ 275

RESULT 4

US-09-052-089A-4
Sequence 4, Application US/09052089A
Patent No. 6346605

GENERAL INFORMATION:
APPLICANT: Lee, Soo Y.

Choi, Yongwon

TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNF RECEPTOR SUPER
FAMILY, AND USES THEREOF

NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.

STREET: 411 Hackensack Ave, Continental Plaza, 4th
Floor

CITY: Hackensack

STATE: New Jersey

COUNTRY: USA

ZIP: 07601

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/052,089A

FILING DATE: 31-Mar-1998

CLASSIFICATION: <Unknown>

ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.

REGISTRATION NUMBER: 26,742

REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1

TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800

TELEFAX: 201-343-1684

INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:

LENGTH: 220 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

HYPOTHETICAL: NO

FRAGMENT TYPE: internal

ORIGINAL SOURCE:
ORGANISM: mouse

SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-052-089A-4

Query Match 85.0%; Score 906; DB 4; Length 220;
Best Local Similarity 86.0%; Pred. No. 3.2e-71;
Matches 185; Conservative 18; Mismatches 12; Indels 0; Gaps 0;

QY 1 RTIINKLFEDLAQEEENVLDRFLKNELDNRYAQLSQDKERDSQVIITDLRDTLEERN 60
Db 1 KTIINKLFEDLAQEEENVLDAEFLKNELDVYKAQLSQDKERDSQAIITDLRDTLEERN 60
QY 61 ATVVSLOALKAEMLCSTLKKOMKYLEQOODETKQAQDEAGRLRSKKKTMQIEILLQS 120
Db 61 ATVESLOALKAEMLCSTLKKOMKYLEQOODETKQAQDEAGRLRSKKKTMQIEILLQS 120
QY 121 QLPVEEEMIRDMGVGQSAVEQLAVYCVSLKREYENLKREARRASGVADKLKRDLFSSRSK 180
Db 121 QREVEEMIRDMGVGQSAVEQLAVYCVSLKREYENLKREARRATGELADRLKRDLYVSSRSK 180

QY 181 LQTVSELDQAKLELSAQKDLQSAKDEKIMSLKK 215
Db 181 LKTLNTELDQAKLELSAQKDLQSAKDEKIMSLKK 215

RESULT 5

US-09-052-089A-2
Sequence 2, Application US/09052089A
Patent No. 6346605

GENERAL INFORMATION:
APPLICANT: Lee, Soo Y.

Choi, Yongwon

TITLE OF INVENTION: SIGNAL TRANSDUCER FOR THE TNF RECEPTOR SUPER
FAMILY, AND USES THEREOF

NUMBER OF SEQUENCES: 16

CORRESPONDENCE ADDRESS:
ADDRESSEE: David A. Jackson, Esq.

STREET: 411 Hackensack Ave, Continental Plaza, 4th
Floor

CITY: Hackensack

STATE: New Jersey

COUNTRY: USA

ZIP: 07601

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/052,089A

FILING DATE: 31-Mar-1998

CLASSIFICATION: <Unknown>

ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.

REGISTRATION NUMBER: 26,742

REFERENCE/DOCKET NUMBER: 600-1-198 CIP 1

TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800

TELEFAX: 201-343-1684

INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:

LENGTH: 470 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

HYPOTHETICAL: NO

FRAGMENT TYPE: <Unknown>

ORIGINAL SOURCE:
ORGANISM: mouse

SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-052-089A-2

Query Match 85.0%; Score 906; DB 4; Length 470;
Best Local Similarity 86.0%; Pred. No. 8.2e-71;
Matches 185; Conservative 18; Mismatches 12; Indels 0; Gaps 0;

QY 1 RTIINKLFEDLAQEEENVLDRFLKNELDNRYAQLSQDKERDSQVIITDLRDTLEERN 60
Db 56 KTIINKLFEDLAQEEENVLDAEFLKNELDVYKAQLSQDKERDSQAIITDLRDTLEERN 115
QY 61 ATVVSLOALKAEMLCSTLKKOMKYLEQOODETKQAQDEAGRLRSKKKTMQIEILLQS 120
Db 116 ATVESLOALKAEMLCSTLKKOMKYLEQOODETKQAQDEAGRLRSKKKTMQIEILLQS 175
QY 121 QLPVEEEMIRDMGVGQSAVEQLAVYCVSLKREYENLKREARRASGVADKLKRDLFSSRSK 180
Db 176 QREVEEMIRDMGVGQSAVEQLAVYCVSLKREYENLKREARRATGELADRLKRDLYVSSRSK 235
QY 181 LQTVSELDQAKLELSAQKDLQSAKDEKIMSLKK 215
Db 236 LKTLNTELDQAKLELSAQKDLQSAKDEKIMSLKK 270


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; GENERAL INFORMATION:
; APPLICANT: Yen, Timothy J.
; TITLE OF INVENTION: Nucleic Acid Encoding a Transiently
; TITLE OF INVENTION: Expressed Kinechochore Protein, and Methods of Use
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Dann, Dorfman, Herrell and Skillman
; STREET: 1601 Market Street Suite 720
; CITY: Philadelphia
; STATE: PA
; COUNTRY: USA
; ZIP: 19103-2307
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US95/16216
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/353,700
; FILING DATE: 09-DEC-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Reed, Janet E.
; REGISTRATION NUMBER: 36,252
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (215) 563-4100
; TELEFAX: (215) 563-4044
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 3248 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
; ANTI-SENSE: NO
; PCT-US95-16216-1

Query Match          14.6%; Score 156; DB 5; Length 3248;
Best Local Similarity 23.6%; Pred. No. 6e-05;
Matches 57; Conservative 55; Mismatches 90; Indels 40; Gaps 7;

QY 19 LDREFLNELDNVRAQLSQDKERKDSQVITLRLDLEERNATVVSLOALGKA-EMLC 77
DB 2299 LDVTLSEKENLTKQIOEKQGLSELDKILSFKSLLEKEQAIEIQKESKTAVENTLQ 2358
QY 78 STLKQ-----MKYLEQOD---ETKQAOEEAGRLRSKMKKTMEQIELLIQS 120
DB 2359 NQLELNEVAVALGODEIMKTEQSDPPLTEEHQULNSIEKLRARLEADEKKQLCVLQ 2418
QY 121 QLEP-----VEEMIRDMGVGSAVEQLAVYCVSLKEKENLKEARKASGEVADK 169
DB 2419 QLKSEHHAHLKGRVENLELEIARTNOHAHLAEENSGEVEYTLKAKTEGQTSLRG 2478
QY 170 LRKDLFSRSKLTQVYSELDQ---AKLEL--KSAQKDLQSADEKIMSLKK---LTM 218
DB 2479 LELDVVIRSEKENLTKELQEKGERISLELINSFENIIQEKQKQVMEKSGSTAMEM 2538
QY 219 LQ 220
DB 2539 LQ 2540

RESULT 9
US-08-795-475-6
; Sequence 6, Application US/08795475
; Patent No. 5965390
; GENERAL INFORMATION:
```

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; APPLICANT: Bjvrck, Lars
; APPLICANT: Sjvbring, Ulf
; TITLE OF INVENTION: PROTEIN L AND HYBRID PROTEINS THEREOF
; NUMBER OF SEQUENCES: 14
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SEED and BERRY LLP
; STREET: 6300 Columbia Center, 701 Fifth Avenue
; CITY: Seattle
; STATE: Washington
; COUNTRY: USA
; ZIP: 98104-7092
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/795,475
; FILING DATE: 11-FEB-1997
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Mcmasters, David D.
; REGISTRATION NUMBER: 33,963
; REFERENCE/DOCKET NUMBER: 100084.402D1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 622-4900
; TELEFAX: (206) 682-6031
; INFORMATION FOR SEQ ID NO: 6:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 443 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-795-475-6

Query Match          13.9%; Score 148.5; DB 2; Length 443;
Best Local Similarity 23.5%; Pred. No. 2.2e-05;
Matches 51; Conservative 56; Mismatches 97; Indels 11; Gaps 6;

QY 9 FDIAQEEENVLDREFLNELDNVRAQLSQK-DKEKRDQVITLRLDLEERNATVVSLO 67
DB 78 YDLAKESST-WDRQRLKELEKEKKEALELIDQASROYHRATALEKELEKKKALYLALD 136
QY 68 QALKAELGCTLKKQKYLEQOODETK---QAQEEAGRLRSKMK--TMDQIELLIQSQ 121
DB 137 QA-SQDYNRANVALEKELETITREQENRNLLGNKLELDQLSSEKEDLTTEKALEEEKQ 195
QY 122 LPEV--EEMIRDMGVGSAVEQLAVYCVSLKEKENLKEARKASGEVADKLRKDLFSRS 179
DB 196 ISDASROSRLRDLASREAKKQVEKDLANLTAELDKYKEDKQISDASRQRLRDLASRE 255
QY 180 KIQTVYSELDAQLEKLSAQKDLQSADEKIMSLKKL 216
DB 256 AKQVEKDLANLTAELDKYKEDKQISDASRQRLRDL 292

RESULT 10
PCT-US93-03077-3
; Sequence 3, Application PC/TUS9303077
; GENERAL INFORMATION:
; APPLICANT: Board of Regents, The University of Texas System
; APPLICANT: Gaynor, Richard B.
; APPLICANT: Wu, Foon Kin
; TITLE OF INVENTION: PROTEIN CELLULAR FACTOR USEFUL FOR
; TITLE OF INVENTION: REGULATING GENE EXPRESSION
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Arnold White & Durkee
; STREET: P.O. Box 4433
; CITY: Houston
; STATE: Texas
; COUNTRY: USA
```


PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/07/862,025
FILING DATE: April 2, 1992
ATTORNEY/AGENT INFORMATION:
NAME: Kammerer, Patricia A.
REGISTRATION NUMBER: 29,775
REFERENCE/DOCKET NUMBER: UFD270PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 713-787-1540
TELEFAX: 713-749-2679
TELEX:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 1093 amino acids
TYPE: AMINO ACID
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
PCT-US93-03077-1

Query Match 13.5%; Score 144; DB 5; Length 1093;
Best Local Similarity 23.8%; Pred. No. 0.00017;
Matches 60; Conservative 47; Mismatches 77; Indels 68; Gaps 10;

QY 11 LAOEENVLDREPLKNELDNVRAOLSQDKERDSQVITDRLPTL---EERNATVVSLO 67
DB 441 LSEKEDVKTYEFLNEKREKREAOILSKERALLIEAFDNLKDEMFYKESSISSLK 500
QY 68 -----QALGKAEMLC---STLKQMKVLEQ-----QODETKAOOE 100
DB 501 DETQRIAEAEKKVQLACKERDAKKEIKNIKELATRLNSSETADLKEKDEQIRGLME 560
QY 101 AG-----RLRSKMTMEQIELLOSQLEPEVEEMIRDMGV---GSAVSQOL 142
DB 561 EGEKLSKQOLNSNIKKLRADKKNENMVAKLNRKVELEBELOHLQVLDGKEVE-- 618
QY 143 AVCYSLKKEYENLKEARKASGEVADKLKDLFSSRSKLTQTVYSELDOAKLELKSQKDL 202
DB 619 -----KQHENIKKL-----NSWVERQEKDL-----GRLOYMDLEKEKN---RSIQAL 660
QY 203 QSADEKMSLRK 214
DB 661 DSAYKELFTDLK 672

RESULT 13
US-09-085-199B-5
Sequence 5, Application US/09085199B
Patent No. 6235879
GENERAL INFORMATION:
APPLICANT: Hayden, Michael R.
APPLICANT: Hackam, Abigail
APPLICANT: Hug, A.H.M. Mahbubul
APPLICANT: Chopra, Vikramjit Singh
APPLICANT: Kalchman, Michael
TITLE OF INVENTION: Apoptosis Modulators That Interact with the
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Opedahl & Larson
STREET: PO Box 5270
CITY: Frisco
STATE: CO
COUNTRY: USA
ZIP: 80443-5270
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Kb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS DOS 5.0
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/085,199B

FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Larson, Marina T.
REGISTRATION NUMBER: 32038
REFERENCE/DOCKET NUMBER: OBC-P-013052
TELECOMMUNICATION INFORMATION:
TELEPHONE: (970) 668-2050
TELEFAX: (970) 668-2052
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 1090
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: no
ORIGINAL SOURCE:
ORGANISM: human
FEATURE:
OTHER INFORMATION: Huntington-interacting protein
US-09-085-199B-5

Query Match 13.3%; Score 142; DB 4; Length 1090;
Best Local Similarity 21.9%; Pred. No. 0.00025;
Matches 53; Conservative 59; Mismatches 90; Indels 40; Gaps 8;

QY 4 INKLFEDLAOEENVLDREPLKNELDNVRAOLSQDKERDSQVITDRLPTLEERNATV 63
DB 421 VNK-----DEKDHILIER--LYREISGLKADL---ENKKTESORVYIOLKGVSELEADL 469
QY 64 VSLQALGKAEMLCSTLKQMKVLEQODETKAOOEAGRLRSKMTMEQIELLOSQLP 123
DB 470 AEQHLRQQAADDEFFLAELDELRLRQREDEKAKRSISTEIERKAQANEQRYSKLEKYS 529
QY 124 EV---EEMIRDMGVGOSAVQLAVCYSLKKEYENLKEA-----RRASG--EYAD 168
DB 530 ELVQNHADLLKRNKNEVTQVSMARQAVDLREKKELEDLSLERSIDGCRKTQEQLEVL 589
QY 169 KLKRDLESSRSKLTQTVSELDOAK-----LELKSQKDLQSA---DKETMSLRK 214
DB 590 SLKQELATSOEQLVLOGLSFTSASQSEANMAAEFALEKREKRDLSVSGAAREEELSLRK 649
QY 215 KL 216
DB 650 EL 651

RESULT 14
US-09-085-199B-4
Sequence 4, Application US/09085199B
Patent No. 6235879
GENERAL INFORMATION:
APPLICANT: Hayden, Michael R.
APPLICANT: Hackam, Abigail
APPLICANT: Hug, A.H.M. Mahbubul
APPLICANT: Chopra, Vikramjit Singh
APPLICANT: Kalchman, Michael
TITLE OF INVENTION: Apoptosis Modulators That Interact with the
NUMBER OF SEQUENCES: 44
CORRESPONDENCE ADDRESS:
ADDRESSEE: Opedahl & Larson
STREET: PO Box 5270
CITY: Frisco
STATE: CO
COUNTRY: USA
ZIP: 80443-5270
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Kb storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: MS DOS 5.0
SOFTWARE: WordPerfect

;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/09/085,199B
;; FILING DATE:
;; CLASSIFICATION:
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Larson, Marina T.
;; REGISTRATION NUMBER: 32038
;; REFERENCE/DOCKET NUMBER: UBC.P-013052
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (970) 668-2050
;; TELEFAX: (970) 668-2052
;; INFORMATION FOR SEQ ID NO: 4:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 914
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: protein
;; HYPOTHETICAL: no
;; ORIGINAL SOURCE:
;; ORGANISM: human
;; FEATURE: Huntington-interacting protein
;; US-09-085-199B-4

Query Match 13.2%; Score 141; DB 4; Length 914;

Best Local Similarity 21.9%; Pred. No. 0.00024;

Matches 53; Conservative 59; Mismatches 90; Indels 40; Gaps 8;

QY 4 INKLFDAEENVLDEFLKNELDNVRAQLSOKDEKRSQVITDRLPTLEERNATV 63
DB 245 VNK-----DEKDHLEI--LYREISGLKADL---ENKKTSSORVVLQKGVSELEADL 293
QY 64 VSLQALGKAEMLSTLKKQKYLEQOODETKQAQEBAGRLRSKMTMEQIELLOSQ 123
DB 294 AEQHLRQADADCEFLAEDELRLQREDTEKAKRSLSIEKRAQANEQKSKLEKYS 353
QY 124 EV---EEMIDMGVGSQSAVEQLAVYCVSLKKEYENLKEA-----RKASG--EYAD 168
DB 354 ELVONHADLRKNAEVTQKQVSMARQAVDLREKKELEDSLEISDQGRTOEQLEYLE 413
QY 169 KLRDLPSSRSKLTQTVYSELQAK-----LELKSQKDLQSA---DKELMSLKK 214
DB 414 SLKQELGTSQRELOVLOGSLTSAQSEANMAAEFALEKEREKDSLVGAHREELSLARK 473
QY 215 KL 216
DB 474 EL 475

RESULT 15

US-09-085-199B-9

Sequence 9, Application US/09085199B

Patent No. 6235879

GENERAL INFORMATION:

APPLICANT: Hayden, Michael R.

APPLICANT: Hackam, Abigail

APPLICANT: Hug, A.H.M. Mahbubul

APPLICANT: Chopra, Vikramjit Singh

APPLICANT: Kalichman, Michael

TITLE OF INVENTION: Apoptosis Modulators That Interact with the

TITLE OF INVENTION: Huntington's Disease Gene

NUMBER OF SEQUENCES: 44

CORRESPONDENCE ADDRESS:

ADDRESSEE: Opedahl & Larson

STREET: PO Box 5270

CITY: Frisco

STATE: CO

COUNTRY: USA

ZIP: 80443-5270

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Kb storage

COMPUTER: IBM compatible

OPERATING SYSTEM: MS DOS 5.0

;; SOFTWARE: WordPerfect
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/09/085,199B
;; FILING DATE:
;; CLASSIFICATION:
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Larson, Marina T.
;; REGISTRATION NUMBER: 32038
;; REFERENCE/DOCKET NUMBER: UBC.P-013052
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (970) 668-2050
;; TELEFAX: (970) 668-2052
;; INFORMATION FOR SEQ ID NO: 9:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 756
;; TYPE: amino acid
;; TOPOLOGY: linear
;; MOLECULE TYPE: protein
;; HYPOTHETICAL: no
;; ORIGINAL SOURCE:
;; ORGANISM: mouse
;; FEATURE:
;; OTHER INFORMATION: Huntington-interacting protein
;; US-09-085-199B-9

Query Match 13.1%; Score 140; DB 4; Length 756;

Best Local Similarity 23.6%; Pred. No. 0.00023;

Matches 60; Conservative 57; Mismatches 81; Indels 56; Gaps 12;

QY 4 INKLFDAEENVLDEFLKNELDNVRAQLSOKDEKRSQVITDRLPTLEERNATV 63
DB 87 VNK-----DEKDHLEI--LYREISGLTGL---DNKKIESQAMQLKGRVSELEAEL 135
QY 64 VSLQALGKAEML-ICSTLKKQKYLEQOODETKQAQEBAGRLRSKMTMEQIELLOSQ 122
DB 136 AE--OQHIGRAMDCEFLRTELDELKRLQREDTEKAKRSLSIEKRAQANEQKSKLEKY 194
QY 123 PEV---EEMIDMGVGSQSAVEQLAVYCVSLKKEYENLKEA-----RKA 162
DB 195 SELVONHADLRKNAEVTQKQVSMARQAVDLREKKELEDSLEISDQGRTOEQLEYLE 247
QY 163 --SGEVAQKLRDLPSSRSKLTQTVYSELQAK-----LELKSQKDLQSA---D 206
DB 248 QEQODVLENLKHELATSNQELQVLSNLETSAQSEAKMLQIATLEKEQSLATVAAGRE 307
QY 207 KEIMSLKKKLTMLQ 220
DB 308 EELSLARQDLESTQ 321

RESULT 16

US-08-533-306A-2

Sequence 2, Application US/08533306A

Patent No. 5837457

GENERAL INFORMATION:

APPLICANT: Liu, Pu

APPLICANT: Collins, Francis S.

APPLICANT: Siciliano, Michael J.

TITLE OF INVENTION: Markers for Detection of Chromosome 16

TITLE OF INVENTION: Rearrangements

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: Harness, Dickey & Pierce, P.L.C.

STREET: P.O. Box 828

CITY: Bloomfield Hills

STATE: MI

COUNTRY: USA

ZIP: 48303

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible


```

: TELEFAX: (810) 641-0270
: INFORMATION FOR SEQ ID NO: 6
: SEQUENCE CHARACTERISTICS:
:   LENGTH: 816 amino acids
:   TYPE: amino acid
:   TOPOLOGY: linear
:   MOLECULE TYPE: protein
: US-08-742-923A-6

```

Query Match	13.0%	Score 139	DB 2:	Length 816:
Best Local Similarity	22.6%	Pred. No. 0.00031:		
Matches	56;	Conservative	92;	Indels 48;
		Mismatches	92;	Gaps 7

QY	10	DLAAEEBVA	-----	LDREFLNELCUNVAQUSQKREKRDQV	11	LDLDFLDE	58
Db	566	DLMOQDELAABERAKROADLKESELBAE	LLSSLGRA	LOADEKERRLEAR	LAQLEEELE	625	
QY	59	RNATVVSU	-----	QOALGKAEMICSTLKQMKYLLQOODE	DTKQAOEAGRLRSKMKTT	112	
Db	626	EOGNEMAMSDVRATQOAEOLSNELATER	STAOKNESAROOLE	ROMKELSLKHEMGA	685		
QY	113	-----	QIHELLQOLPEVEEMIDMGQSAVOLA	VVCYSLKEVYENKE	-----	158	
Db	686	VKSFKSTIALALEAKINGLEQVE	-----	QBARQKQA	TKSLQKQKXKLKEILLQYODE	735	
QY	159	-----	ARKASGEVADKLRDLESSRSKLO	TVVSELDQALBELKSAQDQSDKE	208		
Db	740	KKMAEQKKEQAEKGNARY	-KOLKQOLEBAEESORINANNRRK	LORELDDEATENSNEANGRE	795		
QY	209	IMSLKKLU	216				
Db	799	VNALSKLU	806				

RESULT 20
 US-08-533-306A-4
 Sequence 4, Application US/08533306A
 Patent No. 5837457
 GENERAL INFORMATION:
 APPLICANT: Liu, Pu
 APPLICANT: Collins, Francis S.
 APPLICANT: Siciliano, Michael J.
 APPLICANT: Claxton, David
 TITLE OF INVENTION: Markers for Detection of Chromosome 16
 NUMBER OF SEQUENCES: 14
 CORRESPONDENCE ADDRESSES:
 ADDRESSEE: Harness, Dickey & Pierce, P.L.C.
 STREET: P.O. Box 828
 CITY: Bloomfield Hills
 STATE: MI
 COUNTRY: USA
 ZIP: 48303
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/533.306A
 FILING DATE: September 25, 1995
 CLASSIFICATION: 435
 ATTORNEY/AGENT INFORMATION:
 NAME: Smith, Deann F.
 REGISTRATION NUMBER: 36683
 REFERENCE/DOCKET NUMBER: 2115-00869COB
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (810) 641-1600
 TELEFAX: (810) 641-0270
 INFORMATION FOR SEQ ID NO: 4:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 885 amino acids

TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-533-306A-4

Query Match 13.0%; Score 139; DB 2; Length 885;
Best Local Similarity 22.6%; Pred. No. 0.00035;
Matches 56; Conservative 52; Mismatches 92; Indels 48; Gaps 7;

QY 10 DLAGEENV-----LDREFLNELDNVRAQLSQDKERDSQVITDITLRLTLEE 58
DB 635 DLWQLQEDLMAAERARKQADLEKEELAEELASSLSGNALQDERKRLRLAIALEELEE 694
QY 59 RNATVSL-----QALKAEMLCSTLKKQMYLEQODETRKQAEAGRLRSKMTKE-- 112
DB 695 EOGMEMSDRVKRAQQAQQLSNELATERSTAKNSARQQLERQNKELRSKLHEMGA 754
QY 113 -----QIELLOSQLEPEVEMIRDMGVSAVEQLAVYCVSLKEVENLKE----- 158
DB 755 VKSKFKSTIALEKIAQLEQVE-----QAREKQAA-TKSLKQDKKLKEILLQVEDE 808
QY 159 -----ARKASGVADKLKRLDFSSRSKLTQTVYSELDQAKLELSAQKDLQSDAKE 208
DB 809 RKMAEQYKEQAEKGNARV-KQLKQLEAEESQRIANRRKIQRLDEATEESNEAMGRE 867
QY 209 IMSLKKL 216
DB 868 VNALKSKL 875

RESULT 21
US-08-742-923A-4
Sequence 4, Application US/08742923A
Patent No. 5869611

GENERAL INFORMATION:
APPLICANT: Liu, Pu
APPLICANT: Collins, Francis S.
APPLICANT: Siciliano, Michael J.
APPLICANT: Claxton, David
TITLE OF INVENTION: Markers for Detection of Chromosome 16
TITLE OF INVENTION: Rearrangements
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: Harness, Dickey & Pierce, P.L.C.
STREET: P.O. Box 828
CITY: Bloomfield Hills
STATE: MI
COUNTRY: USA
ZIP: 48303
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/742,923A
FILING DATE: No. 5869611member 1, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Smith, Dean F.
REGISTRATION NUMBER: 36683
REFERENCE/DOCKET NUMBER: 2115-00869DVC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (810) 641-1600
TELEFAX: (810) 641-0270
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 885 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-742-923A-4

Query Match 13.0%; Score 139; DB 2; Length 885;
Best Local Similarity 22.6%; Pred. No. 0.00035;
Matches 56; Conservative 52; Mismatches 92; Indels 48; Gaps 7;

QY 10 DLAGEENV-----LDREFLNELDNVRAQLSQDKERDSQVITDITLRLTLEE 58
DB 635 DLWQLQEDLMAAERARKQADLEKEELAEELASSLSGNALQDERKRLRLAIALEELEE 694
QY 59 RNATVSL-----QALKAEMLCSTLKKQMYLEQODETRKQAEAGRLRSKMTKE-- 112
DB 695 EOGMEMSDRVKRAQQAQQLSNELATERSTAKNSARQQLERQNKELRSKLHEMGA 754
QY 113 -----QIELLOSQLEPEVEMIRDMGVSAVEQLAVYCVSLKEVENLKE----- 158
DB 755 VKSKFKSTIALEKIAQLEQVE-----QAREKQAA-TKSLKQDKKLKEILLQVEDE 808
QY 159 -----ARKASGVADKLKRLDFSSRSKLTQTVYSELDQAKLELSAQKDLQSDAKE 208
DB 809 RKMAEQYKEQAEKGNARV-KQLKQLEAEESQRIANRRKIQRLDEATEESNEAMGRE 867
QY 209 IMSLKKL 216
DB 868 VNALKSKL 875

RESULT 22
US-09-310-187A-1
Sequence 1, Application US/09310187A
Patent No. 6358751

GENERAL INFORMATION:
APPLICANT: Benichou, Gilles
APPLICANT: Pedoseyeva, Eugenia
TITLE OF INVENTION: Involvement of Autoantigens in Cardiac
FILE REFERENCE: UCSF-090
CURRENT APPLICATION NUMBER: US/09/310,187A
CURRENT FILING DATE: 1999-05-12
NUMBER OF SEQ ID NOS: 3
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 1
LENGTH: 1939
TYPE: PRN
ORGANISM: Homo sapiens
US-09-310-187A-1

Query Match 13.0%; Score 138.5; DB 4; Length 1939;
Best Local Similarity 25.2%; Pred. No. 0.001;
Matches 55; Conservative 38; Mismatches 78; Indels 47; Gaps 7;

QY 24 LKNELDNVRAQLSQDKERDSQVITDITLRLTLEARNATVYSLQALGKAMLCSTLKKQ 83
DB 1289 LARLEKEKALISQLTGKLSYTOQMEDLKROLEEBEKKANALAHQSAHRHCDLLR-- 1346
QY 84 MKYLEQOODETK-----QAEAGRLRSKMT-----MQIELLOSQLEPEV 125
DB 1347 ---EYEEETEAKAEIQRVLKSNSEVAQWRITVEDAIQRIEELAEAKKLAQRLODA 1402
QY 126 EEMTRDMGVSAVEQLAVYCVSLK-----EVENLK---EARKASGEVADKLRLDLF 175
DB 1403 EE-----AVEANANCSSLEKTKHRLQNIETEDLMDVENSNAANAALDKKQKNFD 1452
QY 176 SSRSKLTQTVYSELDQAKLELSAQKDLQSDAKEIMSLK 213
DB 1453 KILAEWQKYYEE---SQSELESSQKEARSJSTELFKIK 1487

RESULT 23
US-08-095-737-2
Sequence 2, Application US/08095737
Patent No. 5487979

LENGTH: 316 amino acids

TELEFAX: (703) 836-2021
; INFORMATION FOR SEQ ID NO: 31:

; SEQUENCE CHARACTERISTICS:

```

; SEQUENCE CHARACTERISTICS
; LENGTH: 386
; TYPE: amino acid
; TOPOLOGY: linear

```

RESULT 30
US-09-572-191-2
; Sequence 2, Application US/09572191

```

; Patent No. 6355466
; GENERAL INFORMATION:
; APPLICANT: Beraud, Christophe
; APPLICANT: Sakowicz, Roman
; APPLICANT: Wood, Kenneth
; TITLE OF INVENTION: No. 63554661 motor proteins and methods for
; FILE REFERENCE: 1017
; CURRENT APPLICATION NUMBER: US/09/572,191
; CURRENT FILING DATE: 2000-05-17
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 1388
; TYPE: PRT
; ORGANISM: Human
; US-09-572-191-2

```

```

Query Match      12.4%; Score 132.5; DB 4; Length 1388;
Best Local Similarity 24.5%; Pred. No. 0.0022;
Matches 48; Conservative 44; Mismatches 65; Indels 39; Gaps 8;

```

```

QY 10 DLAGEENVLDREF-LKNELDNVRAQLSOKDKERDSQVIITDRLPTLEER-----59
DB 1203 NLRLSQLLEKMWLIGOLDIDIRKQKNSDPNHPNQOLKNEQDEESIKELAKSKIYEE 1262
QY 60 ----NATVVSLOALGKAMCSTLKKROMKYLEGOODETKQAQAEAGRLRSKM-----108
DB 1263 MLKKAKDLAEVQALYKKEKCLRMTE---VERTQLESFAFEKQRLKLEMYEER 1319
QY 109 -KTEQIETLLQSLPEVEEMIRDMG---VGQSAVEQLAVYCVSLKKEENKEARKASGE 165
DB 1320 ERTSQEMEMLRK---QVECLAENGKLYGHQNHQKIQYVRLKKEVRL-----AE 1368
QY 166 VADKLK-KDLFSSRSK 180
DB 1369 ETEKLRANVFLKEKK 1384

```

```

RESULT 31
US-08-938-105-3
; Sequence 3, Application US/08938105
; Patent No. 6353151
; GENERAL INFORMATION:
; APPLICANT: Leinwand, Leslie A.
; APPLICANT: Vikstrom, Karen L.
; TITLE OF INVENTION: TRANSGENIC MODEL FOR HEART FAILURE
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Sheridan Ross P.C.
; STREET: 1700 Lincoln St., Suite 3500
; CITY: Denver
; STATE: CO
; COUNTRY: U.S.A.
; ZIP: 80203
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/938,105
; FILING DATE:
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Crook, Wanneil M.
; REGISTRATION NUMBER: 31,071
; REFERENCE/DOCKET NUMBER: 3595-4
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (303) 863-9700
; TELEFAX: (303) 863-0223
; INFORMATION FOR SEQ ID NO: 3:

```

```

; SEQUENCE CHARACTERISTICS:
; LENGTH: 1886 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-938-105-3

```

```

Query Match      12.4%; Score 132.5; DB 4; Length 1886;
Best Local Similarity 21.1%; Pred. No. 0.0033;
Matches 64; Conservative 59; Mismatches 80; Indels 101; Gaps 12;

```

```

QY 1 RTINKLFFDLAGEENVLDREF-----EFLK-----25
DB 1078 RAKVEKLRSDLTRELEISERLAEAGATSVQIEMNKKREAFQKMRDLAEATLQHEAT 1137
QY 26 -----NELDNVRAQLSOKDKERDSQVIITDRLPTLEERNATVVSLOQA 69
DB 1138 AALRRKHADSVAEELGIDIDILQRYKQKLEKESEFLKLELDVTSNMQ-----1186
QY 70 LGRKEMLCSTLKKOMKYLEGOODETKQAQAEAGR-----LRSKMT-----MEQI 114
DB 1187 IIRAK---ANLEKVSRTLEDOANEYRYKLEPAQSLNDFTTQRAKLQTEGCELARQLEEK 1243
QY 115 ELLI-----OSQLPEVEEMIRDMVGQSAVE---QLAVY-CVSLKKEYNLKEAR 160
DB 1244 EALIMQLTGRKSLSTQGMEDLKRQLEEGKAKNALAHQLSARHDCDLRQYEEEMK 1303
QY 161 KASEVADKLKRLDFFSSRSKIQT-----VYSELQAKLELSAQKDLOSADKEISLKKL 216
DB 1304 AELQRLVSKANSEVAQWRTKYETDAIQRTTELEBAKKL--AQR-LQDAEAVAVNAK 1360
QY 217 TMLQ 220
DB 1361 SLE 1364

```

```

RESULT 32
US-08-466-390-4
; Sequence 4, Application US/08466390
; Patent No. 5686562
; GENERAL INFORMATION:
; APPLICANT: TOURKATLY, GARY
; APPLICANT: LIDGARD, GRAHAM P
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: TESTA, HURWITZ & THIBEAULT
; STREET: 125 HIGH STREET
; CITY: BOSTON
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/466,390
; FILING DATE: 06-JUN-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: PITCHER ESQ, EDWARD R
; REGISTRATION NUMBER: 27,829
; REFERENCE/DOCKET NUMBER: MTP-013
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 248-7100
; TELEFAX: (617) 248-7100
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2101 amino acids

```

```

;      TYPE: amino acid
;      TOPOLOGY: linear
;      MOLECULE TYPE: protein
US-08-466-390-4

```

Query Match	12.2%	Score 130;	DB 1;	length 2101;
Best Local Similarity	24.9%	Pred. No. 0.0061;		
Matches	57;	Conservative	47;	Mismatches 81;
			Indels	44;
			Gaps	10;

Qy	24	LKNELDNVRAQLSOKDE----	KRDSOVIIDTLEDTEENRAIVSLQALGAEMCST	79
Db	503	LTSLETLNATIQOQDDELGLKQQAKEKQAOLQOTLQOQBOASQGRH---	QVEQLSSS	559
Qy	80	LK--KOMKYLEOQDDETKQ-----	AOEAGRLSRMKMTWEOIELL-----	117
Db	560	LKQKEOOLKEVAKEQOATRDHQAQQLATAABEBSRALREBDALIKQJEALEKEAKLEI	61.9	
Qy	118	LQSOLEPEVEEIBRMGCQSAVEBOLAVNYCYSLKKEYNL-----	KEARKKASGEYADK	169
Db	620	LQOOL-QVANEARDS--AQTSVTAQREKAKELSRKVELELACVETARQOHEAOQAVAE-	675	
Qy	170	LKRLDFSRSKTL--QVIVSELDQAKLELSAQKDLOLQADKELMSLKK	215	
Db	676	LELDLRBQOKATEKERYAOEKDLOLQJQOLAKESLKVYVGSLEEKRR	724	

RESULT 33
US-08-470-950-4
; Sequence 4, Application US/08470950
; Patent No. 5698439

```

1  APPLICANT:  TOURKATY, GARY
2  APPLICANT:  LIDGARD, GRAHAM P
3  TITLE OF INVENTION:  NOVEL MALIGNANT CELL TYPE MARKERS OF THE
4  NUMBER OF INVENTION:  INTERIOR NUCLEAR MATRIX
5  TITLE OF SEQUENCES:  6
6  CORRESPONDENCE ADDRESS:
7  ADDRESSEE:  TESTA, HURWITZ & THIBEAULT
8  STREET:  125 HIGH STREET
9  CITY:  BOSTON
10 STATE:  MA
11 COUNTRY:  USA
12 ZIP:  02110
13
14 COMPUTER READABLE FORM:
15 MEDIUM TYPE:  Floppy disk
16 COMPUTER:  IBM PC compatible
17 OPERATING SYSTEM:  PC-DOS/MS-DOS
18 SOFTWARE:  PatentIn Release #1.0, Version #1.25
19
20 CURRENT APPLICATION DATA:
21 APPLICATION NUMBER:  US/08/470,950
22 FILING DATE:  06-JUN-1995
23 CLASSIFICATION:  536
24
25 ATTORNEY/AGENT INFORMATION:
26 NAME:  PITCHER ESO, EDWARD R
27 REGISTRATION NUMBER:  27,829
28 REFERENCE/DOCKET NUMBER:  MTP-013
29
30 TELECOMMUNICATION INFORMATION:
31 TELEPHONE:  (617) 248-7000
32 TELEFAX:  (617) 248-7100
33
34 INFORMATION FOR SEQ ID NO:  4:
35 SEQUENCE CHARACTERISTICS:
36     LENGTH:  2101 amino acids
37     TYPE:  amino acid
38     TOPOLOGY:  linear
39
40 MOLECULE TYPE:  protein
41
42 LS-08-470-950-4

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Query Match	12.2%	Score 130;	DB 1;	length 2101;
Best Local Similarity	24.9%	Pred. No. 0.0061;		
Matches 57; Conservative	47;	Mismatches 81;	Indels 44;	Gaps 10

[illegible]

RESULT 34
US-08-467-781-4
; Sequence 4, Application US/08467781
; Patent No. 5780596

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Query Match	12.2%	Score 130	DB 1	Length 2101
Best Local	24.9%	Pred. No. 0.0061		
Matches 57	Conservative 47	Mismatches 81	Indels 44	Gaps 10

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Qy 24 LKNEIDNVRADLSOKKRE----KRD5OYIIDLTDLTEERNATVYSLOQALGKRAEMCST 79
    | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db 503 LTSELTTLTNATITGQDDDELAKGKOAKEKQAOLQTLQDQEQASGGRH---QVEQLSSS 559
    | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |

Qy 80 LK--KOMKUYEQOODETKQ-----AOEAGRLRSKMKTWEQIELL----- 117
    | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |
Db 560 LKQKEQDQKEVAGKQKQNTROHQAQQLTAAEBRNASJREBDALQJQLEALEKKAQKLEI 619
    | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |

Qy 118 LOSQLEPEVEEMIRMGSGQASAVEDQAVYCSLKKKEYENL-----KKAQKASGEYADK 169
    | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : | : |

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Db 620 LQOOL-QVANEARDS--AOTSVTQAREKALSRKVEELQACVETAREOHEAOAYAE- 675
QY 170 LKRDLFSSRSKL---QTVYSELDOAKLELSAOKDLOSADKEIMSLKK 215
Db 676 LEIQLRSEQOKATEKERVNAOEKQDLOLQALKEISLKVTKGSLFEERK 724

RESULT 35
US-08-195-487-4
; Sequence 4, Application US/08195487
; Patent No. 5783403
; GENERAL INFORMATION:
; APPLICANT: TOUKATLY, GARY
; APPLICANT: LIDGARD, GRAHAM P
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: TESTA HURWITZ & THIBEAULT
; STREET: 53 STATE STREET
; CITY: BOSTON
; STATE: MA
; COUNTRY: USA
; ZIP: 02109
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/195,487
; FILING DATE:
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/07/901,701
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: PITCHER ESO, EDMUND R
; REGISTRATION NUMBER: 27,829
; REFERENCE/DOCKET NUMBER: MTP-013
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617/248-7000
; TELEFAX: 617/248-7100
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2101 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-195-487-4

Query Match 12.2%; Score 130; DB 1; Length 2101;
Best Local Similarity 24.9%; Pred. No. 0.0061;
Matches 57; Conservative 47; Mismatches 81; Indels 44; Gaps 10;

QY 24 LKNELDNVRQALSQKDE---KRDQVYIIDTLTLEERNATVYSLQALGRAEMLCST 79
Db 503 LTSLETLTNMTIQOQDELGLKQAKQAKQALQTLQOOEQASQGLRH---QVEQLSSS 559
QY 80 LK---KOMKYLEQOQDETKQ-----AOEAGRLRSKMKKTMEQIELL----- 117
Db 560 LKQKEQQLKEVAEKQEXTRODHAAQOLATTAEEERASLRERDAALQLEALEKEKAALLET 619
QY 118 LQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKREYENL-----KEARKASGEVADK 169
Db 620 LQOOL-QVANEARDS--AOTSVTQAREKALSRKVEELQACVETAREOHEAOAYAE- 675
QY 170 LKRDLFSSRSKL---QTVYSELDOAKLELSAOKDLOSADKEIMSLKK 215
Db 676 LEIQLRSEQOKATEKERVNAOEKQDLOLQALKEISLKVTKGSLFEERK 724

RESULT 36
US-08-483-924-4
; Sequence 4, Application US/08483924
; Patent No. 5882876
; GENERAL INFORMATION:
; APPLICANT: TOUKATLY, GARY
; APPLICANT: LIDGARD, GRAHAM P
; TITLE OF INVENTION: NOVEL MALIGNANT CELL TYPE MARKERS OF THE
; TITLE OF INVENTION: INTERIOR NUCLEAR MATRIX
; NUMBER OF SEQUENCES: 6
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: TESTA, HURWITZ & THIBEAULT
; STREET: 125 HIGH STREET
; CITY: BOSTON
; STATE: MA
; COUNTRY: USA
; ZIP: 02110
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/483,924
; FILING DATE: 07-JUN-1995
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: PITCHER ESO, EDMUND R
; REGISTRATION NUMBER: 27,829
; REFERENCE/DOCKET NUMBER: MTP-013
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 248-7000
; TELEFAX: (617) 248-7100
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 2101 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-483-924-4

Query Match 12.2%; Score 130; DB 2; Length 2101;
Best Local Similarity 24.9%; Pred. No. 0.0061;
Matches 57; Conservative 47; Mismatches 81; Indels 44; Gaps 10;

QY 24 LKNELDNVRQALSQKDE---KRDQVYIIDTLTLEERNATVYSLQALGRAEMLCST 79
Db 503 LTSLETLTNMTIQOQDELGLKQAKQAKQALQTLQOOEQASQGLRH---QVEQLSSS 559
QY 80 LK---KOMKYLEQOQDETKQ-----AOEAGRLRSKMKKTMEQIELL----- 117
Db 560 LKQKEQQLKEVAEKQEXTRODHAAQOLATTAEEERASLRERDAALQLEALEKEKAALLET 619
QY 118 LQSQLPEVEEMIRDMGVGQSAVEQLAVYCVSLKREYENL-----KEARKASGEVADK 169
Db 620 LQOOL-QVANEARDS--AOTSVTQAREKALSRKVEELQACVETAREOHEAOAYAE- 675
QY 170 LKRDLFSSRSKL---QTVYSELDOAKLELSAOKDLOSADKEIMSLKK 215
Db 676 LEIQLRSEQOKATEKERVNAOEKQDLOLQALKEISLKVTKGSLFEERK 724

RESULT 37
US-09-452-294-1
; Sequence 1, Application US/09452294
; Patent No. 6287790
; GENERAL INFORMATION:
; APPLICANT: Bissell, Mita
; APPLICANT: Lelievre, Sophie
; TITLE OF INVENTION: UTILIZATION OF NUCLEAR STRUCTURAL PROTEINS FOR TARGETED
; TITLE OF INVENTION: THERAPY AND DETECTION OF PROLIFERATIVE AND
; TITLE OF INVENTION: DIFFERENTIATION DISORDERS

Db 8 FILEQOERKLOQKEL--QIDS---LLOQEKELSSSLHOKLCSFOEEMAKENLFEEL 61
QY 67 00ALGKAEMLCSTLKQKMYLEQOODETKQOEGAGRLRSKMKTME-----QIE 115
Db 62 KQTLDELDKLOQKEQOERLTKQLEEKRSRAEELKLEELKLGKREKELEKSSAHTQAT 121
QY 116 LLLQSOQLPEVEEMIRDMGVGOSAVEOL-AVYCVSLKKEYENLKEARKASGEVADKLKRD 174
Db 122 LLLLEKIDSMVQSLVEDYTAQFESYKALTASIEDLKLNSLQEKVAKAGKADVDQHQI 181
QY 175 FSSRSKLOTVYSELDOAKLELKSQKDLQSDAKEI-MSLKKKLTMLQ 220
Db 182 LATESSNQEVYRML-LDLQTKSALKETEI--KEITVSFLQKITDLO 224

RESULT 40
US-08-700-178-2
; Sequence 2, Application US/08700178
; Patent No. 5783669
; Patent No. 5783669 5700912
; GENERAL INFORMATION:
; APPLICANT: Hawkins, Phillip R.
; APPLICANT: Wilde, Craig G.
; APPLICANT: Sellhammer, Jeffrey J.
; TITLE OF INVENTION: HYALURONAN RECEPTOR EXPRESSED IN HUMAN
; TITLE OF INVENTION: UMBILICAL VEIN ENDOTHELIAL CELLS
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
; STREET: 3174 Porter Drive
; CITY: Palo Alto
; STATE: CA
; COUNTRY: USA
; ZIP: 94304
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/700,178
; FILING DATE: August 20, 1996
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/402,217
; FILING DATE: March 10, 1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Billings, Lucy J.
; REGISTRATION NUMBER: 36,749
; REFERENCE/DOCKET NUMBER: PF-0028-1 DIV
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-855-0555
; TELEFAX: 415-845-4166
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 351 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-700-178-2

Query Match 12.1%; Score 129.5; DB 1; Length 351;
Best Local Similarity 25.6%; Pred. NO. 0.00073;
Matches 58; Conservative 47; Mismatches 97; Indels 25; Gaps 8;

QY 9 FDLAQEEENVLDREFLNKELDNVRAQLSQKDEKRD--QVILDTLRDPLEERNATVSL 66
Db 8 FILEQOERKLOQKEL--QIDS---LLOQEKELSSSLHOKLCSFOEEMAKENLFEEL 61
QY 67 00ALGKAEMLCSTLKQKMYLEQOODETKQOEGAGRLRSKMKTME-----QIE 115
Db 62 KQTLDELDKLOQKEQOERLTKQLEEKRSRAEELKLEELKLGKREKELEKSSAHTQAT 121

QY 116 LLLQSOQLPEVEEMIRDMGVGOSAVEOL-AVYCVSLKKEYENLKEARKASGEVADKLKRD 174
Db 122 LLLLEKIDSMVQSLVEDYTAQFESYKALTASIEDLKLNSLQEKVAKAGKADVDQHQI 181
QY 175 FSSRSKLOTVYSELDOAKLELKSQKDLQSDAKEI-MSLKKKLTMLQ 220
Db 182 LATESSNQEVYRML-LDLQTKSALKETEI--KEITVSFLQKITDLO 224

Search completed: September 4, 2002, 16:11:00
Job time: 7599 sec

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